SEQUENCE LISTING

<110> National Institute of Agrobiological Sciences

<120> Sodium/Proton antiporter gene

<130> MOA-006PCT

<150> JP 1998-365604

<151> 1998-12-22

<160> 2

<170> PatentIn Ver. 2.0

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<212> DNA

<213> Oryza sativa

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<222> (297).. (1901)

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tetgcgaatc gaatictitg tittititic tetaatiita eegggaattg tegaattagg 180
catteaccaa egagcaagag gggagtggat tggttggtta aagcteegea tettgeggeg 240
gaaatetege tetetietet geggtggtg geeggagaag tegeegeegg tgagge atg 299
Met

ggg atg gag gtg gcg gcg cgg ctg ggg gct ctg tac acg acc tcc 347 Gly Met Glu Val Ala Ala Ala Arg Leu Gly Ala Leu Tyr Thr Thr Ser 5 10 15

1

gac tac gcg tcg gtg gtg tcc atc aac ctg ttc gtc gcg ctg ctc tgc 395

Asp Tyr Ala Ser Val Val Ser Ile Asn Leu Phe Val Ala Leu Leu Cys

20 25 30

gcc tgc atc gtc ctc ggc cac ctc ctc gag gag aat cgc tgg gtc aat 443
Ala Cys Ile Val Leu Gly His Leu Leu Glu Glu Asn Arg Trp Val Asn
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gag tcc aic acc gcg ctc atc atc ggg ctc tgc acc ggc gtg gtg atc 491 Glu Ser Ile Thr Ala Leu Ile Ile Gly Leu Cys Thr Gly Val Val Ile

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gat	ctc	ttc	ttc	atc	tac	ctc	ctc	cct	ccg	atc	atc	ttc	aat	gca	ggt	587
Asp	Leu	Phe	Phe	He	Tyr	Leu	Leu	Pro	Pro	Ile	Ile	Phe	Asn	Ala	Gly	
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ttt	cag	gta	aag	aaa	aag	caa	ttc	ttc	cgg	aat	ttc	atg	acg	atc	aca	635
Phe	Gln	Val	Lys	Lys	Lys	Gln	Phe	Phe	Arg	Asn	Phe	Met	Thr	Ile	Thr	
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tta	itt	gga	gcc	gtc	ggg	aca	atg	ata	tcc	ttt	ttc	aca	ata	tct	att	683
Leu	Phe	Gly	Ala	Val	Gly	Thr	Met	Ile	Ser	Phe	Phe	Thr	Ile	Ser	Ile	
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gct	gcc	att	gca	ata	ttc	agc	aga	atg	aac	att	gga	acg	ctg	gat	gta	731
Ala	Ala	Ile	Ala	Ile	Phe	Ser	Arg	Met	Asn	Ile	Gly	Thr	Leu	Asp	Val	
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gga	gat	ttt	ctt	gca	att	gga	gcc	atc	ttt	tct	gcg	aca	gat	tct	gtc	779
Gly	Asp	Phe	Leu	Ala	Ile	Gly	Ala	Ile	Phe	Ser	Ala	Thr	Asp	Ser	Val	
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tgc	aca	t t g	cag	gtc	ctc	aat	cag	gat	gag	aca	ccc	ttt	ttg	tac	agt	827
Cys	Thr	Leu	Gln	Val	Leu	Asn	Gln	Asp	Glu	Thr	Pro	Phe	Leu	Tyr	Ser	
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Leu	Val	Phe	Gly	Glu	Gly	Val	Val	Asn	Asp	Ala	Thr	Ser	Ile	Val	Leu	
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Phe	Asn	Ala	Leu	Gln	Asn	Phe	Asp	Leu	Val	His	Ile	Asp	Ala	Ala	Val	
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ttc	ctt	gga	gta	ttt	gct	gga	ttg	ctc	agt	gca	tac	ata	atc	aag	aag	1019
Phe	Leu	Gly	Val	Phe	Ala	Gly	Leu	Leu	Ser	Ala	Tyr	Ile	Ile	Lys	Lys	
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cta	tac	att	gga	agg	cat	tct	act	gac	cgt	gag	gtt	gcc	ctt	atg	atg	1067
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ctc atg gct tac cit tca tat atg ctg gct gag ttg cta gat tig agc

Leu Met Ala Tyr Leu Ser Tyr Met Leu Ala Glu Leu Leu Asp Leu Ser

1115

260	265	270
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Trp	His	Asn	Val	Thr	Glu	Ser	Ser	Arg	Val	Thr	Thr	Lys	His	Ala	Phe	
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Gly	Lys	Ser	Ile	Gly	Ile	Ser	Ser	Ile	Leu	Leu	Gly	Leu	Val	Leu	Ile	
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ctc ctg aca agc atg caa ggt tct gac ctc gag agt aca acc aac att

Leu Leu Thr Ser Met Gln Gly Ser Asp Leu Glu Ser Thr Thr Asn Ile

į

ı	170	475	480

gtg agg cct tcc agc ctc cgg atg ctc ctc acc aag ccg acc cac act 1787 Val Arg Pro Ser Ser Leu Arg Met Leu Leu Thr Lys Pro Thr His Thr 490 495 485

1835 gtc cac tac tac tgg cgc aag ttc gac gac gcg ctg atg cga ccg atg Val His Tyr Tyr Trp Arg Lys Phe Asp Asp Ala Leu Met Arg Pro Met 510 505 500

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1931 Gln Ser His Gly Gly Arg 535 530

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2330

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<213> Oryza sativa

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Cys Ala Cys Ile Val Leu Gly His Leu Leu Glu Glu Asn Arg Trp Val

35 40 45

Asn Glu Scr Ile Thr Ala Leu Ile Ile Gly Leu Cys Thr Gly Val Val
50 55 60

Ile Leu Leu Met Thr Lys Gly Lys Ser Ser His Leu Phe Val Phe Ser

Harry Street World Street

Glu Asp Leu Phe Phe Ile Tyr Leu Leu Pro Pro Ile Ile Phe Asn Ala

Gly Phe Gln Val Lys Lys Gln Phe Phe Arg Asn Phe Met Thr Ile

Thr Leu Phe Gly Ala Val Gly Thr Met Ile Ser Phe Phe Thr Ile Scr

Ile Ala Ala Ile Ala Ile Phc Ser Arg Met Asn Ile Gly Thr Leu Asp

Val Gly Asp Phe Leu Ala Ilc Gly Ala Ilc Phe Ser Ala Thr Asp Ser

Val Cys Thr Leu Gln Val Leu Asn Gln Asp Glu Thr Pro Phe Leu Tyr

Ser Leu Val Phe Gly Glu Gly Val Val Asn Asp Ala Thr Ser Ile Val

Leu Phe Asn Ala Leu Gln Asn Phe Asp Leu Val His Ile Asp Ala Ala

Val Val Leu Lys Phe Leu Gly Asn Phe Phe Tyr Leu Phe Leu Scr Ser 210 215 220

Thr Phe Leu Gly Val Phe Ala Gly Leu Leu Ser Ala Tyr Ile Ile Lys
225 230 235 240

Lys Leu Tyr Ile Gly Arg His Ser Thr Asp Arg Glu Val Ala Leu Met
245 250 255

Met Leu Met Ala Tyr Leu Ser Tyr Met Leu Ala Glu Leu Leu Asp Leu 260 265 270

Ser Gly Ile Leu Thr Val Phe Phe Cys Gly Ile Val Mct Ser His Tyr 275 280 285

Thr Trp His Asn Val Thr Glu Ser Ser Arg Val Thr Thr Lys His Ala
290 295 300

Phe Ala Thr Leu Scr Phe Ile Ala Glu Thr Phe Leu Phe Leu Tyr Val
305 310 315 320

Gly Met Asp Ala Leu Asp Ile Glu Lys Trp Glu Phe Ala Ser Asp Arg
325 330 335

Pro Gly Lys Scr Ile Gly Ile Scr Scr Ile Leu Leu Gly Leu Val Leu 340 345 350

Ile Gly Arg Ala Ala Phe Val Phe Pro Leu Ser Phe Leu Ser Asn Leu 355 360 365

Thr Lys Lys Ala Pro Asn Glu Lys Ile Thr Trp Arg Gln Gln Val Val 370 375 380

Ile Trp Trp Ala Gly Leu Met Arg Gly Ala Val Ser Ile Ala Leu Ala 385 390 395 400

Tyr Asn Lys Phe Thr Arg Ser Gly His Thr Gln Leu His Gly Asn Ala
405 410 415

Ile Met Ile Thr Ser Thr Ile Thr Val Val Leu Phe Ser Thr Met Val
420 425 430

Phe Gly Met Met Thr Lys Pro Leu Ile Arg Leu Leu Pro Ala Ser 435 440 445

Gly His Pro Val Thr Ser Glu Pro Ser Ser Pro Lys Ser Leu His Ser 450 455 460

Pro Leu Leu Thr Ser Met Gln Gly Ser Asp Leu Glu Ser Thr Thr Asn 465 470 475 480

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Thr Val His Tyr Tyr Trp Arg Lys Phe Asp Asp Ala Leu Met Arg Pro
500 510

Met Phe Gly Gly Arg Gly Phe Val Pro Phe Ser Pro Gly Ser Pro Thr 515 520 525

Glu Gln Ser His Gly Gly Arg

530 535